

FIGURE 1

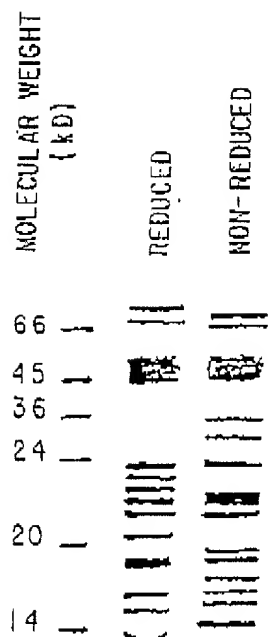


FIGURE ■ 2.

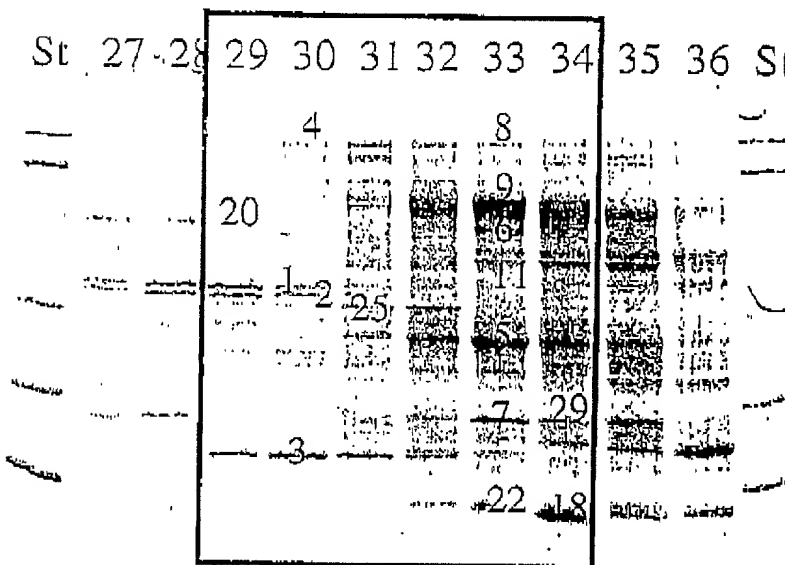
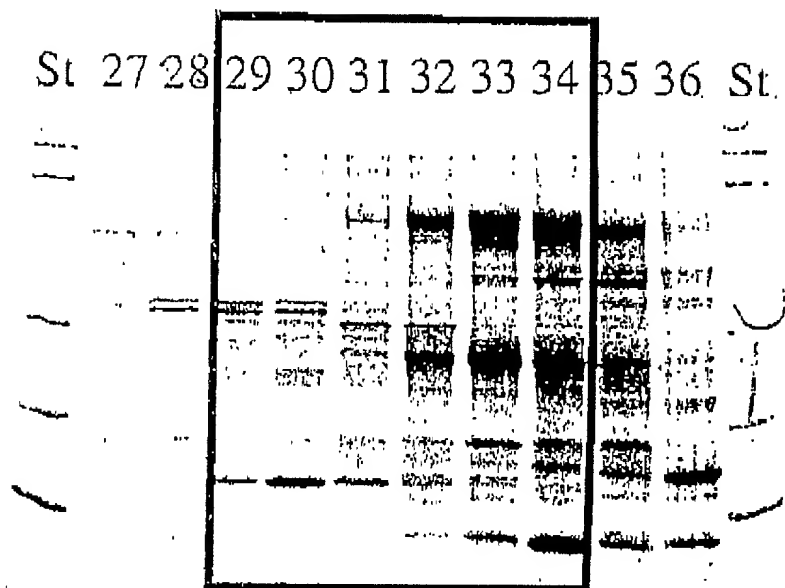
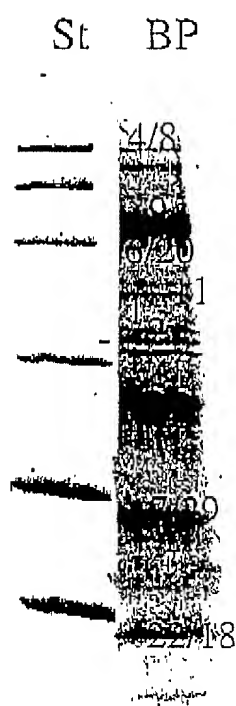


FIGURE ■ 3



Band No.	Identity
1	histone H1.c
2	histone H1.c
3	ribosomal protein RS20
4	similar to ribosomal protein LORP
5	BMP-3
6	$\alpha$ 2 macroglobulin RAP & BMP-3
7	similar to ribosomal protein LORP
8	BMP-3
9	BMP-3
11	ribosomal protein RL6 & BMP-3
18	TGF- $\beta$ 2/SPP24
20	Factor H
22	TGF- $\beta$ 2
25	BMP-3 & H1.x
29	BMP-3 & ribosomal protein RL32

FIGURE 4

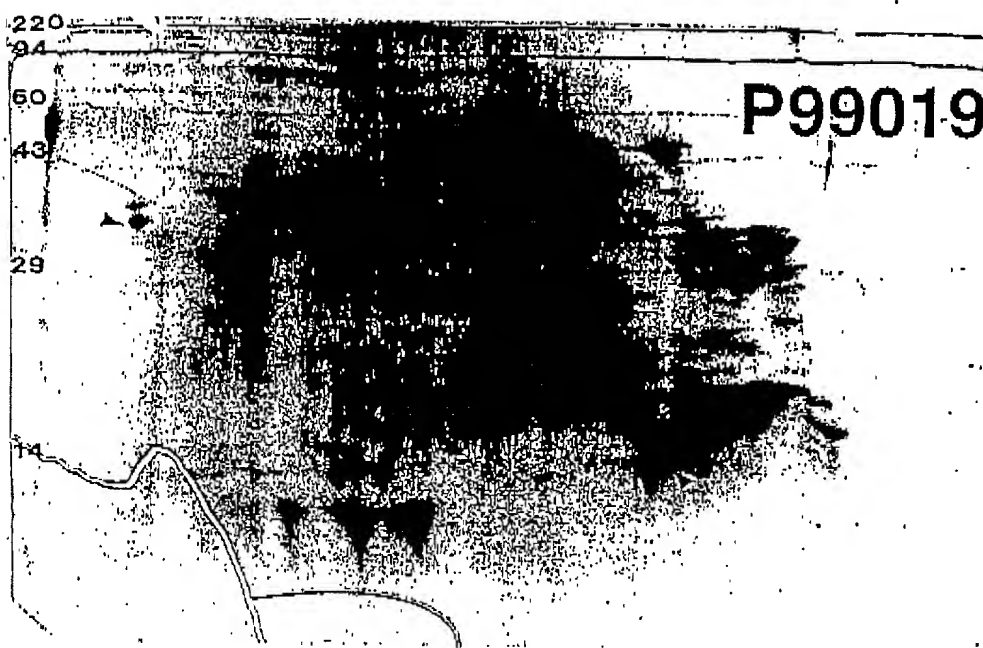
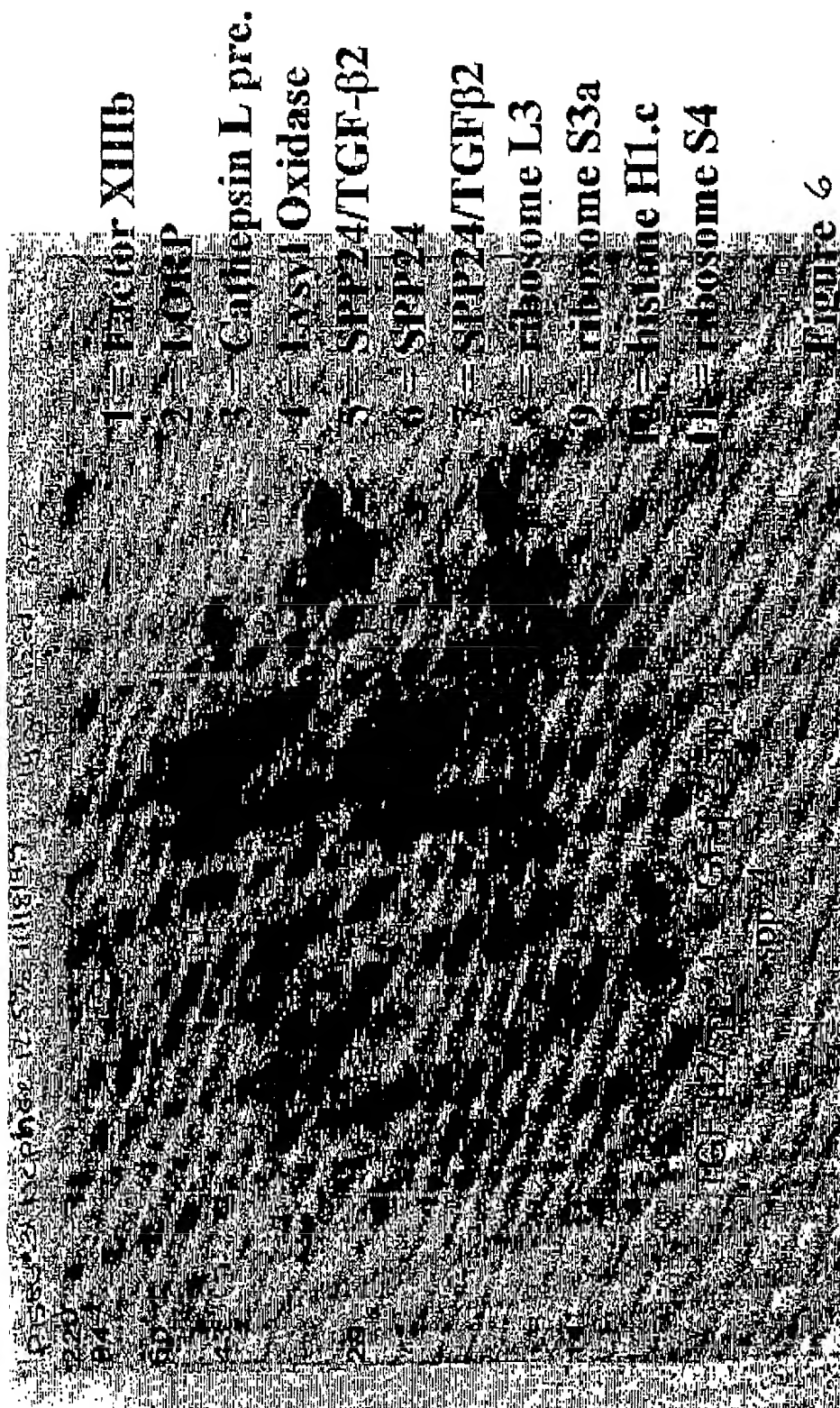


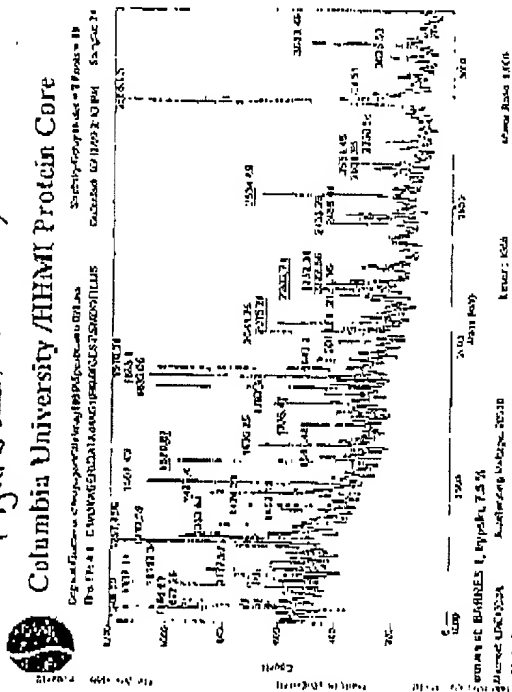
FIGURE 5

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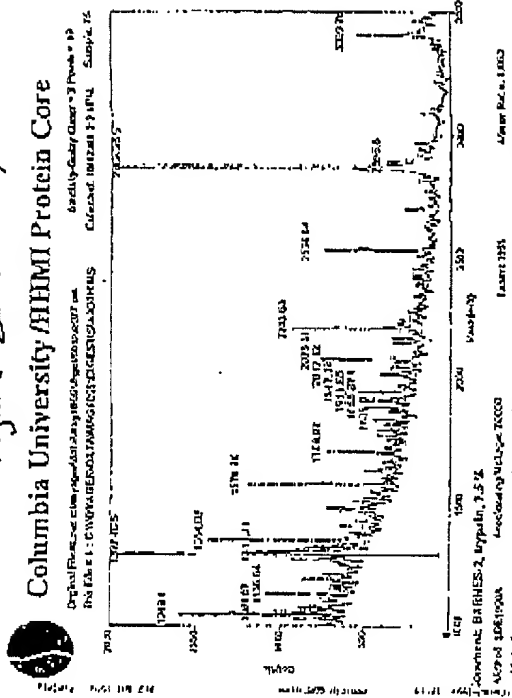


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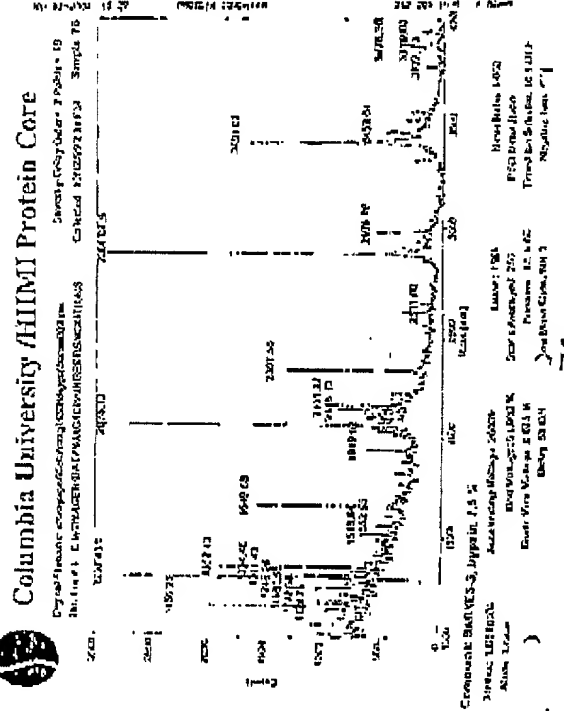
7A  
Figure 7A (Band 1)



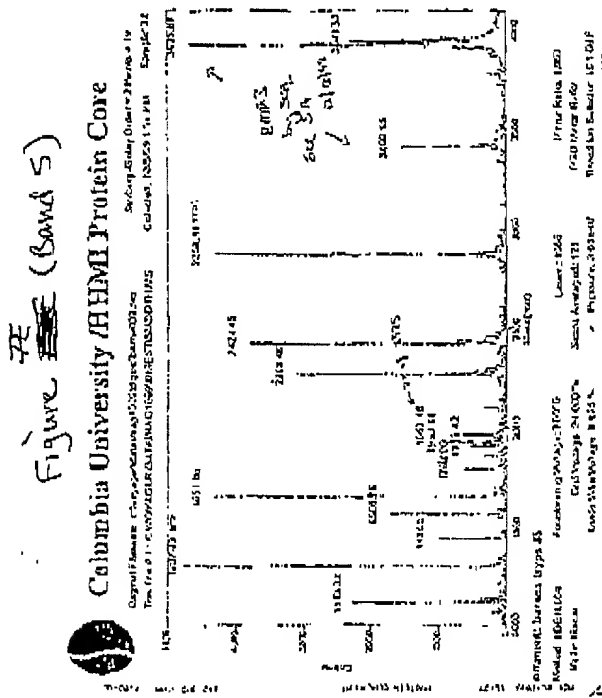
7B  
Figure 7B (Band 2)

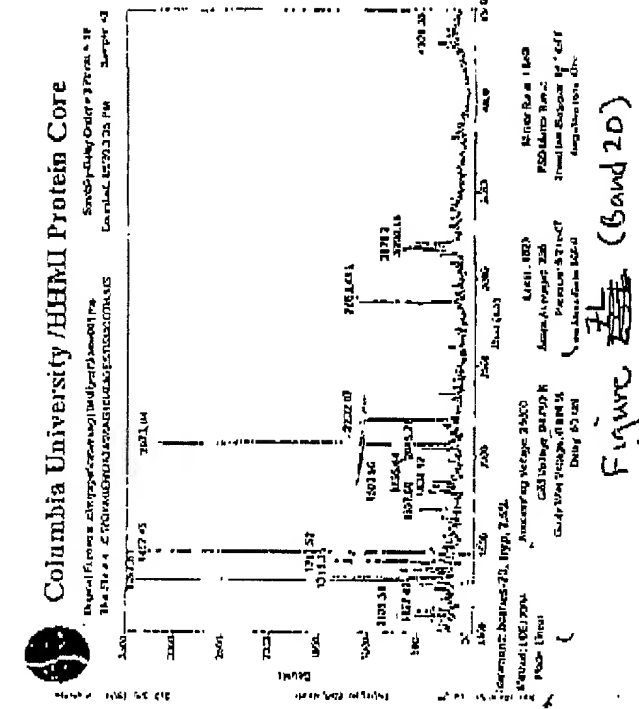
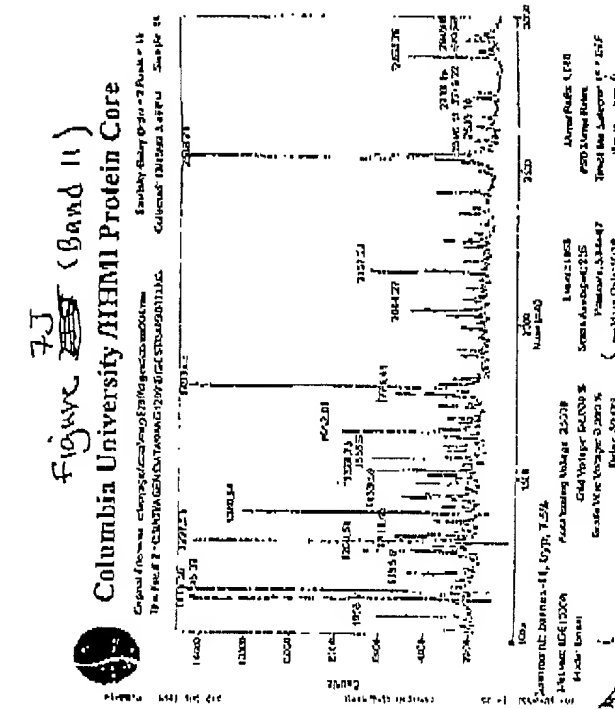
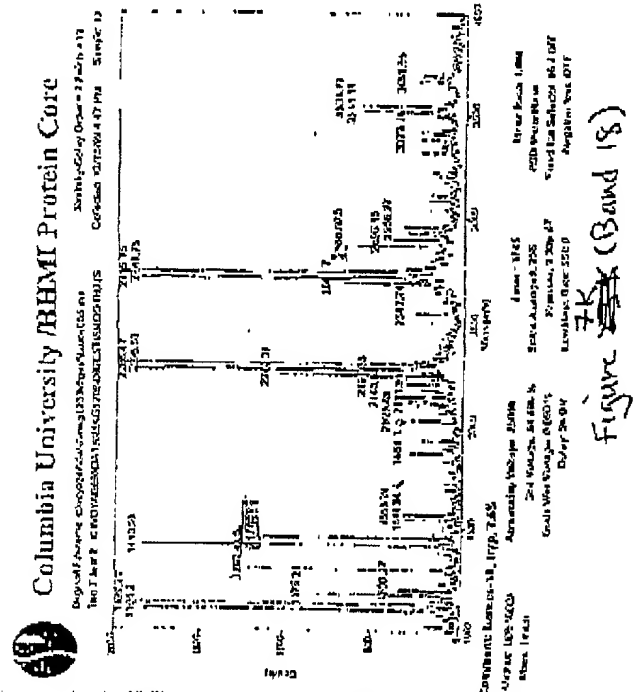
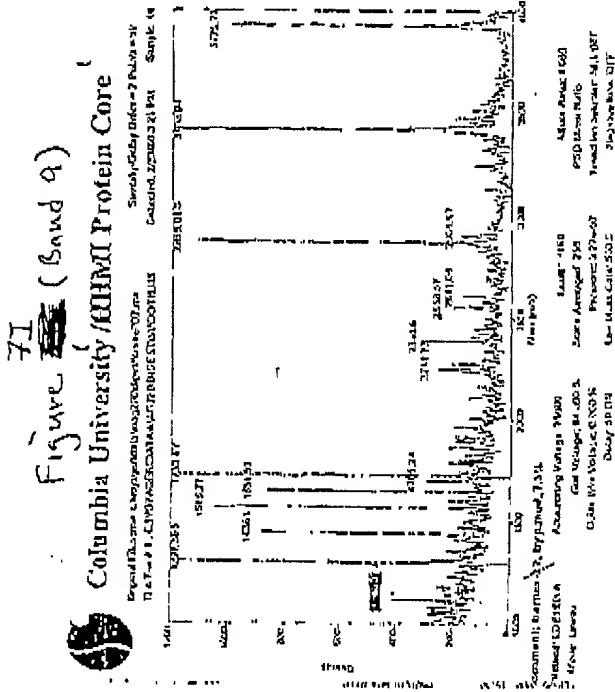


7C  
Figure 7C (Band 3)



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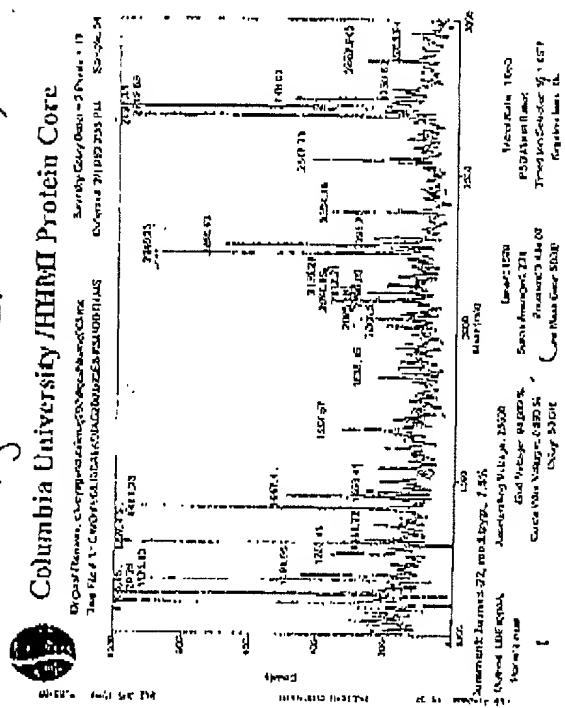




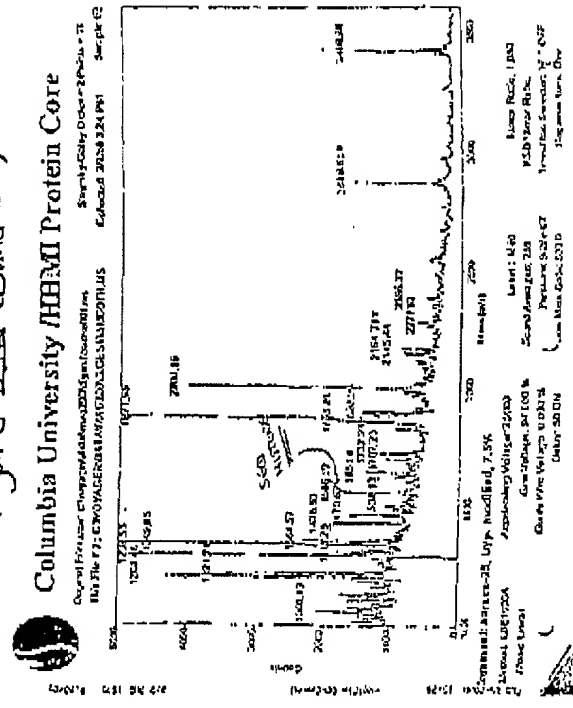


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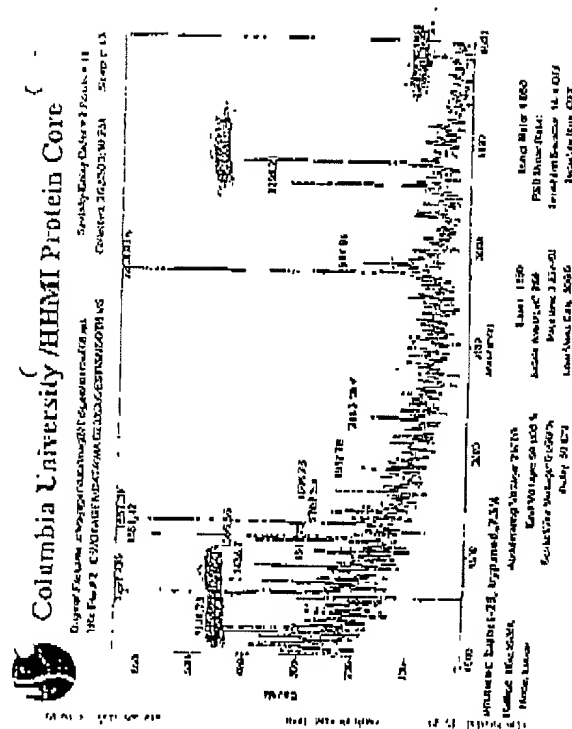
7M Figure 7N (Band 22)



7N Figure 7N (Band 25)



70 Figure 7O (Band 29)



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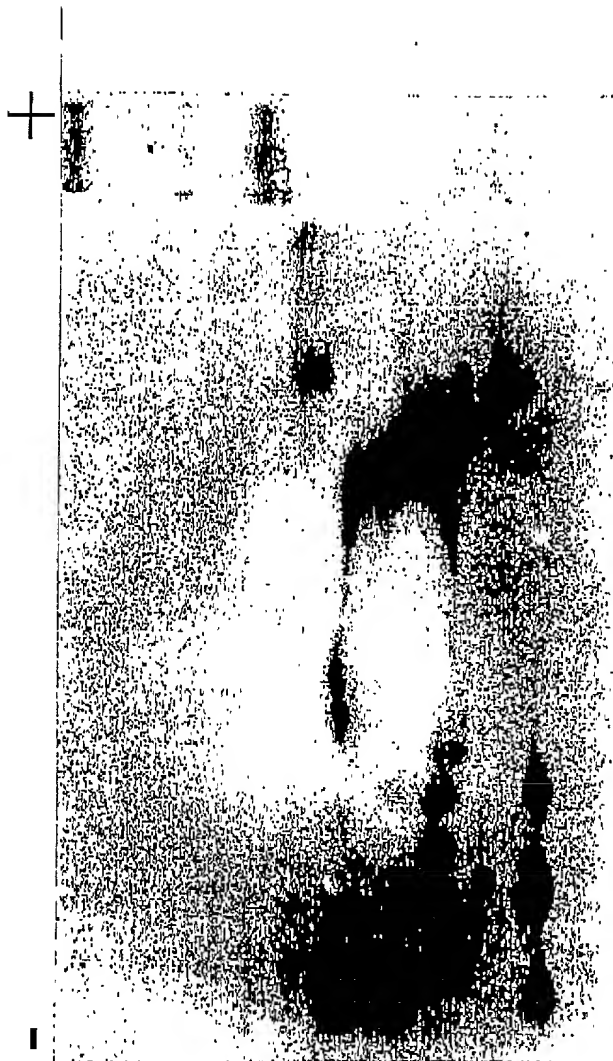


FIGURE 8

h2

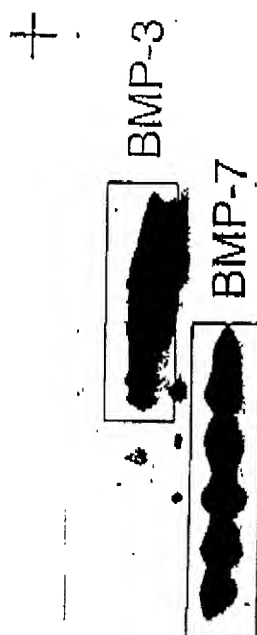


FIGURE 9B

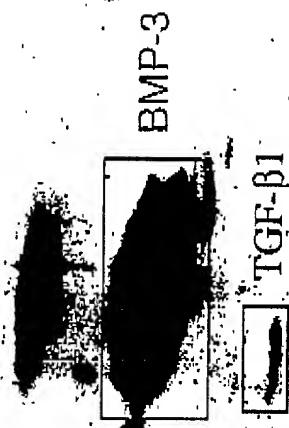


FIGURE 23-9D



FIGURE 9A



FIGURE 15.9C

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FIGURE 10

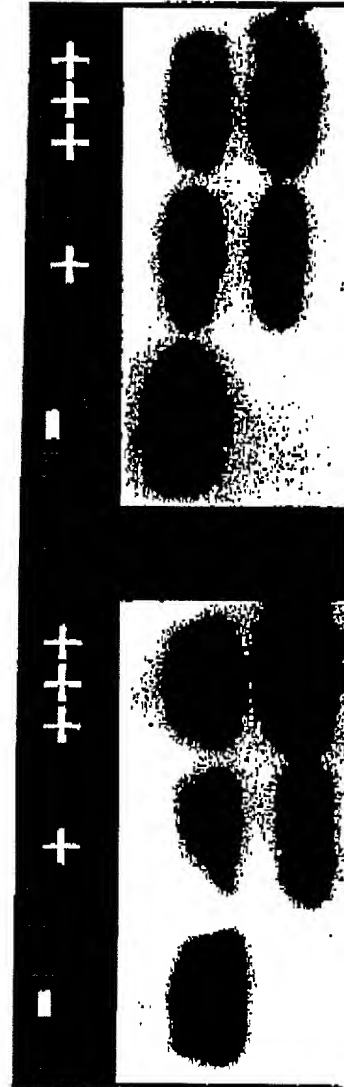
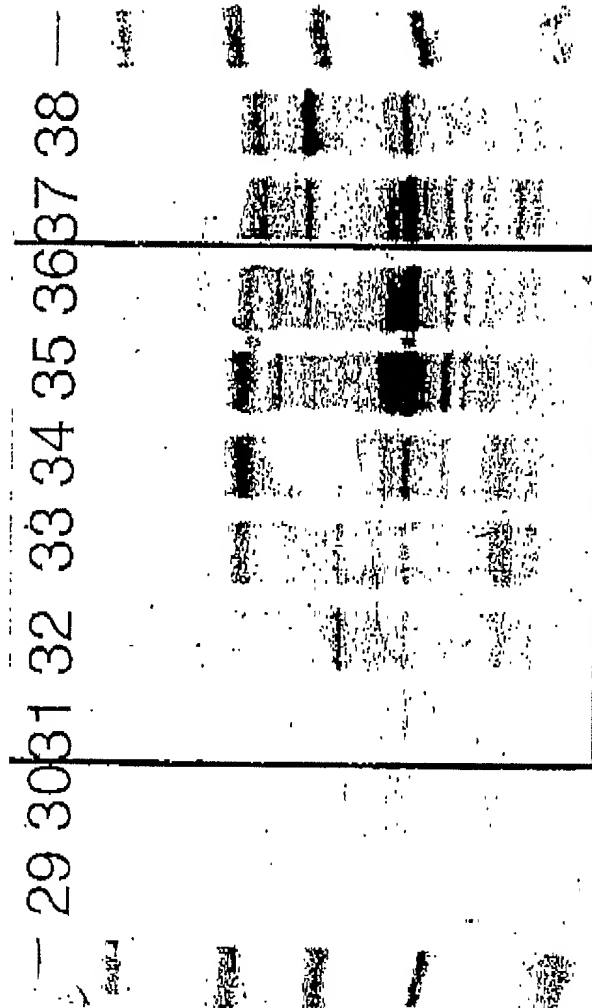


FIGURE 11

FIGURE 12

FIGURE 13A

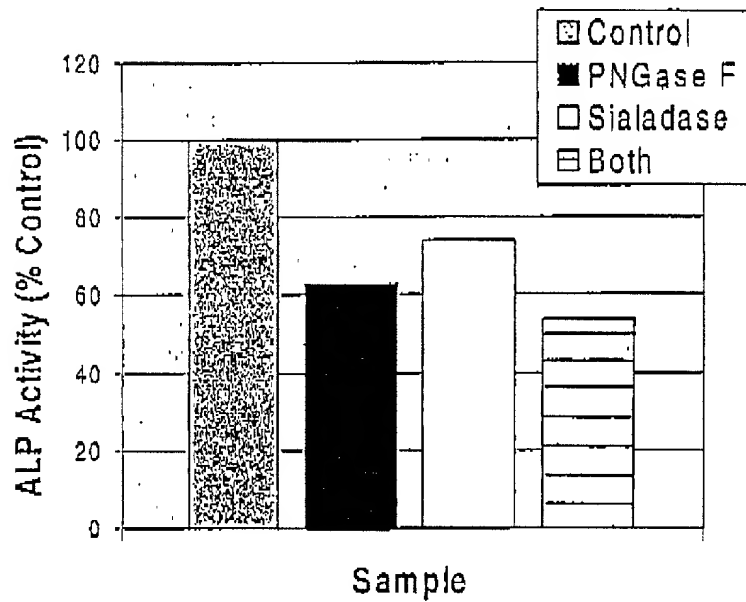
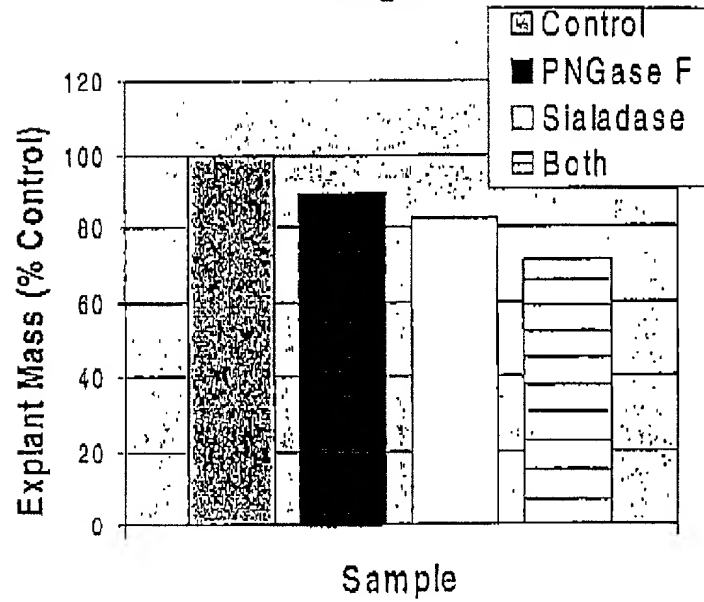


FIGURE 13B

Figure 14 Antibody Listing

Specificity	Antigen	Host Species	PC/MC	Source	Catalog No.
TGF-b1 (human)	Protein	Rabbit	Polyclonal	Promega	G1221
TGF-b2 (human)	Peptide	Rabbit	Polyclonal	Santa Cruz Biotechnology	sc-90
TGF-b3 (human)	Peptide	Rabbit	Polyclonal	Santa Cruz Biotechnology	sc-82
BMP-2 (human)	Protein	Rabbit	Polyclonal	Austral Biologics	PA-513-9
BMP-3 (human)	Peptide	Chicken	Polyclonal	Research Genetics	NA
BMP-4 (human)	Peptide	Goat	Polyclonal	Santa Cruz Biotechnology	sc-6896
BMP-5 (human)	Peptide	Goat	Polyclonal	Santa Cruz Biotechnology	sc-7405
BMP-6 (human)	Peptide	Mouse	Monoclonal	Novocastria Laboratories	NCL-BMP6
BMP-7 (human)	Peptide	Rabbit	Polyclonal	Research Genetics	NA
FGF-1 (human)	Peptide	Goat	Polyclonal	Santa Cruz Biotechnology	sc-1884
osteonectin (bovine)	Protein	Mouse	Monoclonal	DSEB	AON-1
osteocalcin (bovine)	Protein	Rabbit	Polyclonal	Accurate Chemicals	A761/R111
serum albumin (bovine)	Protein	Rabbit	Polyclonal	Chemicon International	AB870
transferrin (human)	Protein	Chicken	Polyclonal	Chemicon International	AB797
apo-A1 lipoprotein (human)	Protein	Goat	Polyclonal	Chemicon International	AB740

Figure 15A Identification of Proteins by Amino Acid Sequencing of Tryptic Fragments from 1D Gels

Band	Sample	Sequence Data	Best Database Match	Match	Identification	Species	Acc. No.	AAs
1								
2	fx 49 (1579)	XLAAAGYDVEK	ALAAAGYDVEK	11/11	histone H1.c	human	87668 (NCBI)	65-75
3	fx 67 (1346)	SLEKVCADLR	SLEKVCADLR	11/11	40s Ribosomal Protein S20	rat	R3RT20 (PIR)	31-41
4	fx 65 ()	(V)VGMLGFPSEAPV	VVGMLGFPSEKRV	11/14	LORP	mouse	AAC95338 (NCBI)	213- 228
5	N terminal seq	STGVLLPLQNNELPG	STGVLLPLQNNELPG	15/15	BMP-3	human	4557371 (NCBI)	290- 304
	fx 72 (3925)	STGVLLPLQNNELPGA EYQY	STGVLLPLQNNELPGA AEYQY	20/20	BMP-3	human	4557371 (NCBI)	290- 309
	fx 74 (3409)	STGVLLPLQ	STGVLLPLQ	9/9	BMP-3	human	4557371 (NCBI)	290- 298
6	fx 55 (1586)	(S)QTLOQFE	SQTLOQFE	7/8	BMP-3	human	4557371 (NCBI)	346- 353
	fx 47	VYAF	no match		777			
	N terminal seq	HAGKYSREKNT(P)A(P )	HGGKYSREKNGPQP	11/14	$\alpha$ 2-Macroglobulin Receptor Assoc. Pro.	human	P30533 (Swiss-Prot)	31-46
	fx 57 (1438)	SQTLOQFEQ	SQTLOQFEQ	9/9	BMP-3	human	4557371 (NCBI)	346- 354
	fx 57 (1852)	SLKPSNHA	SLKPSNHA	8/8	BMP-3	human	4557371 (NCBI)	410- 417
7	fx 51 (1093)	AALRPLVKP	AALRPLVKP	9/9	60s Ribosomal Protein L32	mouse	P17832 (Swiss-Prot)	1-8
	fx 37 (no MS)	A(H)Q)VERYV	AVER	5/5	60s Ribosomal Protein L32	mouse	P17832 (Swiss-Prot)	109- 113
	fx 37 (no MS)	A(H)Q)VERYV	HQSDRYV	5/7	60s Ribosomal Protein L32	mouse	P17832 (Swiss-Prot)	22-28
8	fx 76 ()	XALF(G)AQLGXALGP1	no match		777			
9	fx 58 (1587)	SQTLOQFEQT	SQTLOQFEQT	10/10	BMP-3	human	P12645 (Swiss-Prot)	346- 355

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Figure 15B Identification of Proteins by Amino Acid Sequencing of Tryptic Fragments from 1D Gels

Band	Sample	Sequence Data	Best Database Match	Match	Identification	Species	Acc. No.	AA#
11	fx 55 (1311)	SQTLXF	SQTLQIF	5/6	BMP-3	human	4557371 (NCBI)	348-351
	fx 47 (1772)	VLATVTKPVGGDK	VLATVTKPVGGDK	13/13	60s Ribosomal Protein L6	human	Q02878 (Swiss-Prot)	67-99
	fx 76 (1785)	xVFAL	VFAL	4/4	60s Ribosomal Protein L6	human	Q02878 (Swiss-Prot)	273-276
	fx 51 (1145)	AVPQLQGYLR	AVPQLQGYLR	9/10	60s Ribosomal Protein L6	human	Q02878 (Swiss-Prot)	262-271
18								
22	fx 58 (1107)	ALDAAYCFR	ALDAAYCFR	9/9	TGF- $\beta$ 2	human	P08112 (Swiss-Prot)	303-311
	fx 68 (no match)	GYNANFCAGACPYL	GYNANFCAGACPYL	14/14	TGF- $\beta$ 2	human	P08112 (Swiss-Prot)	340-353
	fx 66 (1411.71)	VNSQSLSPY	VNSQSLSPY	9/9	SP24	bovine	Q27967 (Swiss-Prot)	42-50
25	fx 39 (1470)	KAAPSV(P)	KAAPSV(P)	8/8	Histone H4 x	human	JC4926 (PIR)	199-206
29								

fx = fraction number (molecular weight of fragment, as measured by SDS-PAGE)



Figure 1A Identification of Proteins by Mass Spectrometry of Tryptic Fragments from 1D Gels

Band	Mass Spec Profile	Species	Acc. No.	Mass Spec Data	Mass Spec Database	Mass Difference	AAS	% Coverage	Comments
1	4 peaks match with histone H1.c	human	87668 (NCBI)	1172.97	1172.37	0.60	110-121	22	15 MS peaks match with Band 2
				1579.87	1579.71	0.16	65-79		
				1708.47	1707.89	0.58	64-78		
				2011.58	2012.32	-0.74	35-54		
2	3 peaks match with histone H1.c	human	87668 (NCBI)	1579.76	1579.71	0.05	65-79	16	identification of slamed peptide confirmed by sequence analysis
				1708.02	1707.99	0.13	64-79		
				2012.12	2012.32	-0.20	35-54		
3	7 peaks match with ribosome S20	rat	R3R20 (PIR)	1129.76	1129.40	0.36	50-59	62	15 MS peaks match with Band 1
				1156.21	1156.30	-0.09	76-83		
				1334.46	1334.62	-0.16	58-66		
				1352.13	1351.58	0.55	88-99		
				1518.04	1517.77	0.27	9-21		
				1919.02	1919.19	-0.17	5-21		
				3404.02	3404.87	-0.85	88-119		
4	3 peaks match with Lysyl Oxidase RP	human	NP002309 (Swiss-Prot)	1997.95	1986.27	-0.32	150-167	8	12 MS peaks match with Band 8
				2410.35	2410.63	-0.28	648-668		
				2610.57	2610.10	0.47	455-478		

Figure 16B Identification of Proteins by Mass Spectrometry of Tryptic Fragments from 1D Gels

Band	Mass Spec Profile	Species	Acc. No.	Mass Spec Data	Mass Spec Database	Mass Difference	AAs	% Coverage	Comments
5	9 peaks match with BMP-3	human	4557371 (NCBI)	1113.32	1113.31	0.01	361-368	48	% coverage calculation is relative to the mature BMP-3, 183 AAs (280-472)
				1438.53	1438.58	-0.05	346-357		
				1566.76	1566.76	0.00	345-357		
				1651.88	1651.91	-0.03	410-424		
				1794.09	1794.02	0.07	346-380		
				2288.46	2288.63	-0.17	374-392		
				2424.45	2424.81	-0.36	373-392		
				3408.15	3407.77	1.38	290-318		
				1002.24	1002.15	0.09	283-290		
6	3 peaks match with $\alpha 2$ -Macroglobulin RAP	human	P30533 (Swiss-Prot)	2362.58	2362.43	0.15	129-150	17	
				3048.51	3048.52	-0.01	257-282		
				1566.93	1566.75	0.18	346-357		
				1651.88	1651.91	-0.03	410-424		
	2 peaks match with BMP-3	human	4557371 (NCBI)					15	% coverage calculation is relative to the mature BMP-3, 183 AAs (280-472)



Figure 16D Identification of Proteins by Mass Spectrometry of Trypic Fragments from 1D Gels

Band	Mass Spec Profile	Species	Acc. No.	Mass Spec Data	Mass Spec Database	Mass Difference	AAs	% Coverage	Comments
11	5 peaks match with BMP-3	human	4557371 (NCBI)	1113.23	1113.31	-0.08	361-368	48	% coverage calculation is relative to the mature BMP-3, 183 AAs (290-472)
				1651.73	1651.91	-0.18	410-424		
				1793.58	1794.02	-0.44	346-360		
				2424.24	2424.81	-0.57	373-392		
				3408.34	3407.77	0.57	290-318		
18	5 peaks match with ribosome L6	human	Q02878 (Swiss-Prot)	1140.36	1140.23	0.15	114-122	16	
				1526.88	1526.86	0.02	141-155		
		mouse	P47911 (Swiss-Prot)	1059.15	1059.12	0.03	10-28		
				1145.36	1145.35	0.01	262-271		
				1386.74	1386.88	0.06	260-271		
	4 peaks match with TGF- $\beta$ 2	human	P08172 (Swiss-Prot)	1101.20	1101.26	-0.06	303-311	52	
				1175.26	1175.42	-0.16	400-409		
				2240.37	2240.60	-0.23	312-328		
				2691.70	2691.91	-0.21	340-362		
				1410.93	1411.60	-0.67	42-53		
	5 peaks match with SPP24	bovine	Q27967 (Swiss-Prot)	1447.59	1447.65	-0.06	113-124	30	
				1540.94	1540.60	0.04	86-98		
				1869.10	1869.05	0.05	62-77		
				2268.47	2268.57	-0.10	33-53		

Figure 16E Identification of Proteins by Mass Spectrometry of Tryptic Fragments from 1D Gels

Band	Mass Spec Profile	Species	Acc. No.	Mass Spec Data	Mass Spec Database	Mass Difference	AAS	% Coverage	Comments
22	5 peaks match with TGF- $\beta$ 2	human	P08112 (Swiss-Prot)	1101.15	1101.26	-0.11	303-311	83	
				1175.13	1175.42	-0.29	400-409		
				2084.16	2084.42	-0.26	312-347		
				2240.25	2240.60	-0.35	312-328		
				2691.61	2691.91	-0.30	340-362		
25	2 peaks match with SPP24	bovine	Q27967 (Swiss-Prot)	1411.23	1411.60	-0.37	42-53	11	
				1447.40	1447.65	-0.25	113-124		
				1208.46	1208.40	0.06	48-57	14	
				1221.71	1222.35	-0.64	107-118		
				1349.65	1350.52	-0.67	107-119		
	5 peaks match with histone H1.x	human	JC4826 (PIR)	1394.57	1394.59	-0.02	48-58		
				1732.23	1732.97	-0.74	43-57		
				1060.43	1060.20	0.23	102-111	31	% coverage calculation is relative to the mature BMP-3, 183 AAS (290-472)
				1438.83	1438.58	0.25	348-357		
				1566.92	1566.76	0.16	345-357		
26	5 peaks match with BMP-3	human	4557371 (NCBI)	1651.80	1651.91	-0.11	410-424		
				3408.86	3407.77	1.09	290-318		



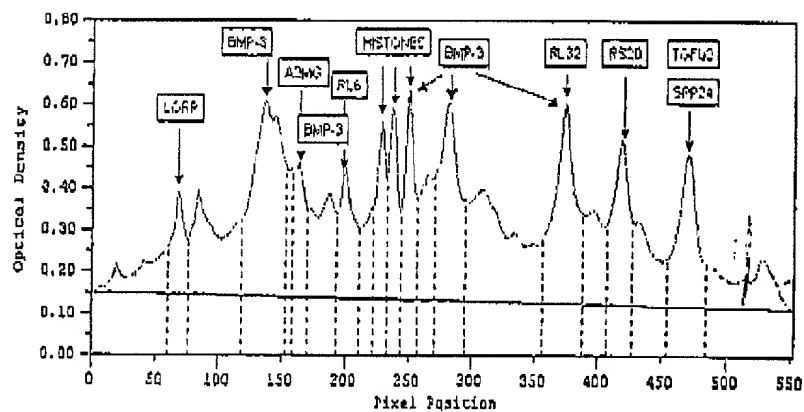


FIGURE 17A



FIGURE 17B

Figure /8 Quantitation of Identified BP proteins

Identified Protein	Percentage of Total Protein
LORP	2
BMP-3	11
BMP - 3 & A2-MG	3
RL6 & BMP-3	4
Histone	3
Histone	3
Histone & BMP-3	4
BMP-3	8
RL32 & BMP-3	8
RS20	5
SPP24 & TGF- $\beta$ 2	6
Total	56%

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Figure 17C Identification of Proteins by Mass Spectrometry of Fragments from 2D Gels

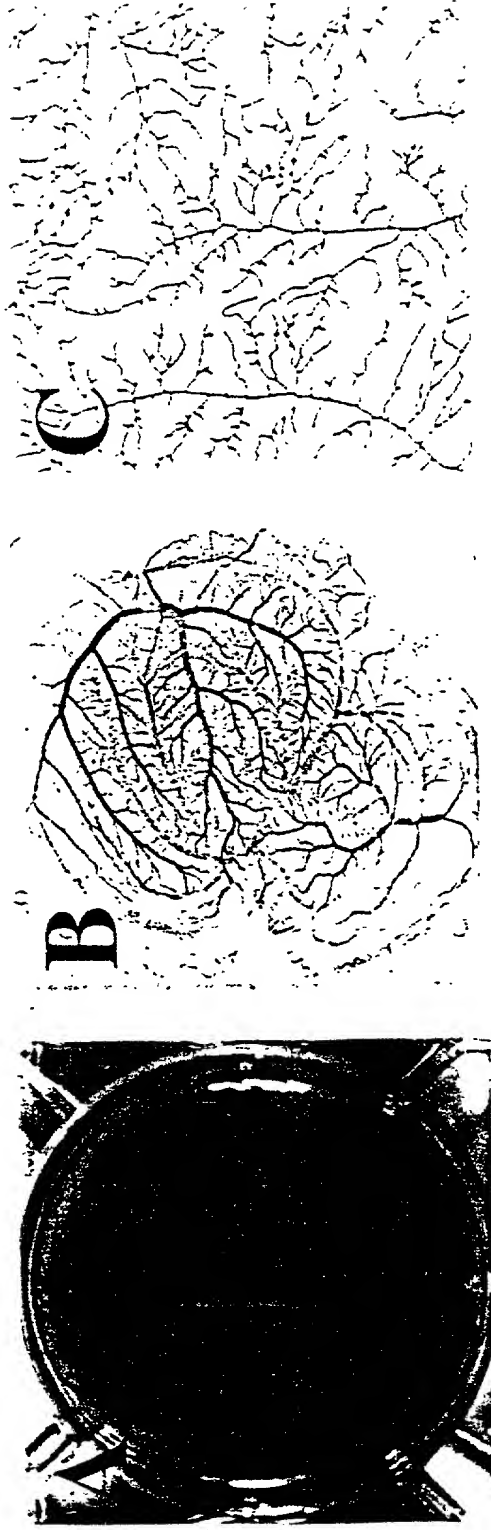
Spot	Digest	Mass Spec Profile	Species	Acc. No.	MS peaks			AAs	% Coverage	Comments
					Data	Database	Diff			
7	Lys-C	4 peaks match with TGF- $\beta$ 2	Bovine	P21214 (Swiss-Prot)	774.56	774.90	-0.34	26-31	42	
					809.69	809.94	-0.25	32-37		
					1175.12	1175.43	-0.31	98-107		
					3168.10	3166.66	1.44	1-25		
					2187.77	2187.51	0.26	42-60	10	
8	Trypsin	12 peaks match with ribosome L3	Bovine	P39872 (Swiss-Prot)	917.39	917.14	0.25	348-355	37	
					984.23	984.15	0.08	10-18		
					1192.62	1192.40	0.22	286-296		
					1380.67	1380.65	0.02	249-260		
					1464.60	1464.63	0.17	103-114		
					1620.86	1620.82	0.04	103-115		
					1778.04	1778.00	-0.16	34-49		
					2238.43	2238.55	-0.12	30-49		
					2325.99	2325.65	0.34	177-197		
					2661.31	2661.04	0.27	200-223		
					2897.94	2898.43	-0.49	70-98		
					2946.10	2948.35	-0.25	198-223		

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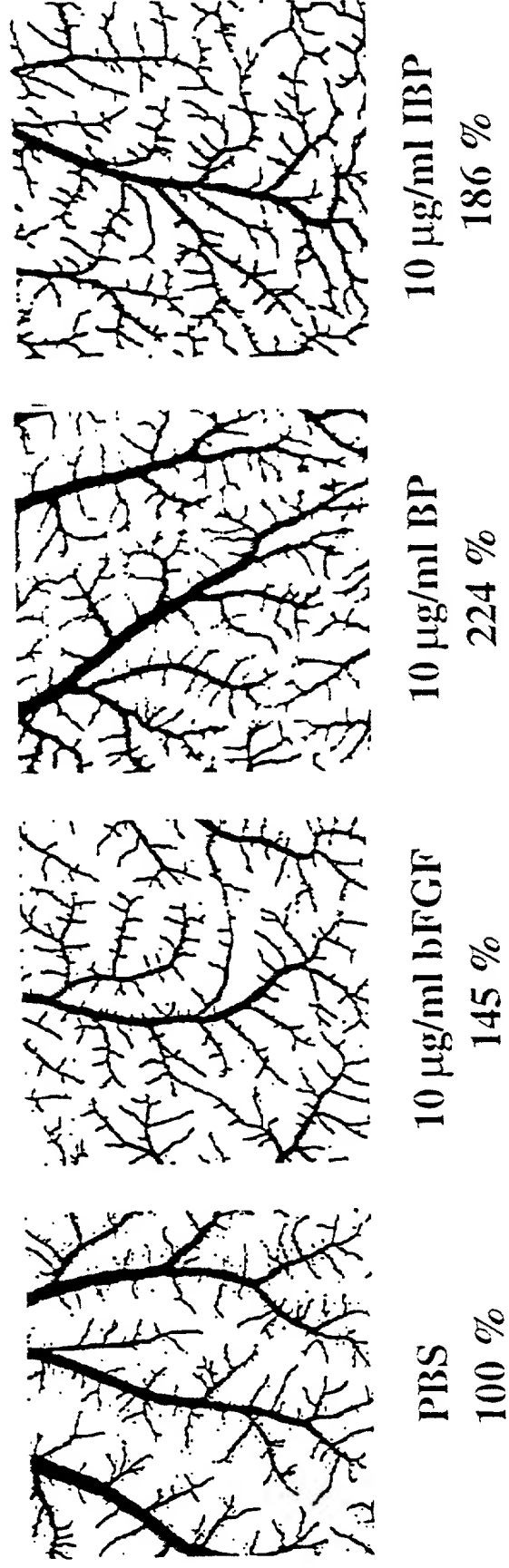
Figure 7B Identification of Proteins by Mass Spectrometry of Fragments from 2D Gels

Spot	Digest	Mass Spec Profile	Species	Acc. No.	MS Peaks			AAs	% Coverage	Comments
					Data	Database	Diff			
9	Trypsin	7 peaks match with ribosome S3a	Mouse	P97351 (Swiss-Prot)	920.05	920.10	-0.05	19-26	28	
					1218.29	1218.31	-0.02	152-161		
					1346.62	1346.49	0.13	151-161		
					1516.69	1516.69	0.00	174-185		
					1583.72	1583.82	-0.10	94-108		
					1719.91	1720.00	-0.09	199-212		
					1953.12	1953.16	-0.04	65-81		
10	Trypsin	4 peaks match with histone H1.c	Human	87558 (NCBI)	1327.75	1327.56	0.19	34-46	23	
					1579.70	1579.71	-0.01	65-79		
					1707.65	1707.89	-0.24	64-79		
					2147.17	2147.53	-0.36	1-21		
					1868.48	1868.38	0.10	230-239	23	
11	Trypsin	6 peaks match with ribosome S4	Human	P12750 (Swiss-Prot)	1216.38	1216.39	0.00	134-144		
					1354.03	1353.61	0.42	230-241		
					1507.81	1507.89	0.12	198-210		
					1557.75	1557.96	-0.23	37-48		
					2140.34	2140.58	-0.24	221-239		
					2591.80	2591.90	-0.10	77-99		

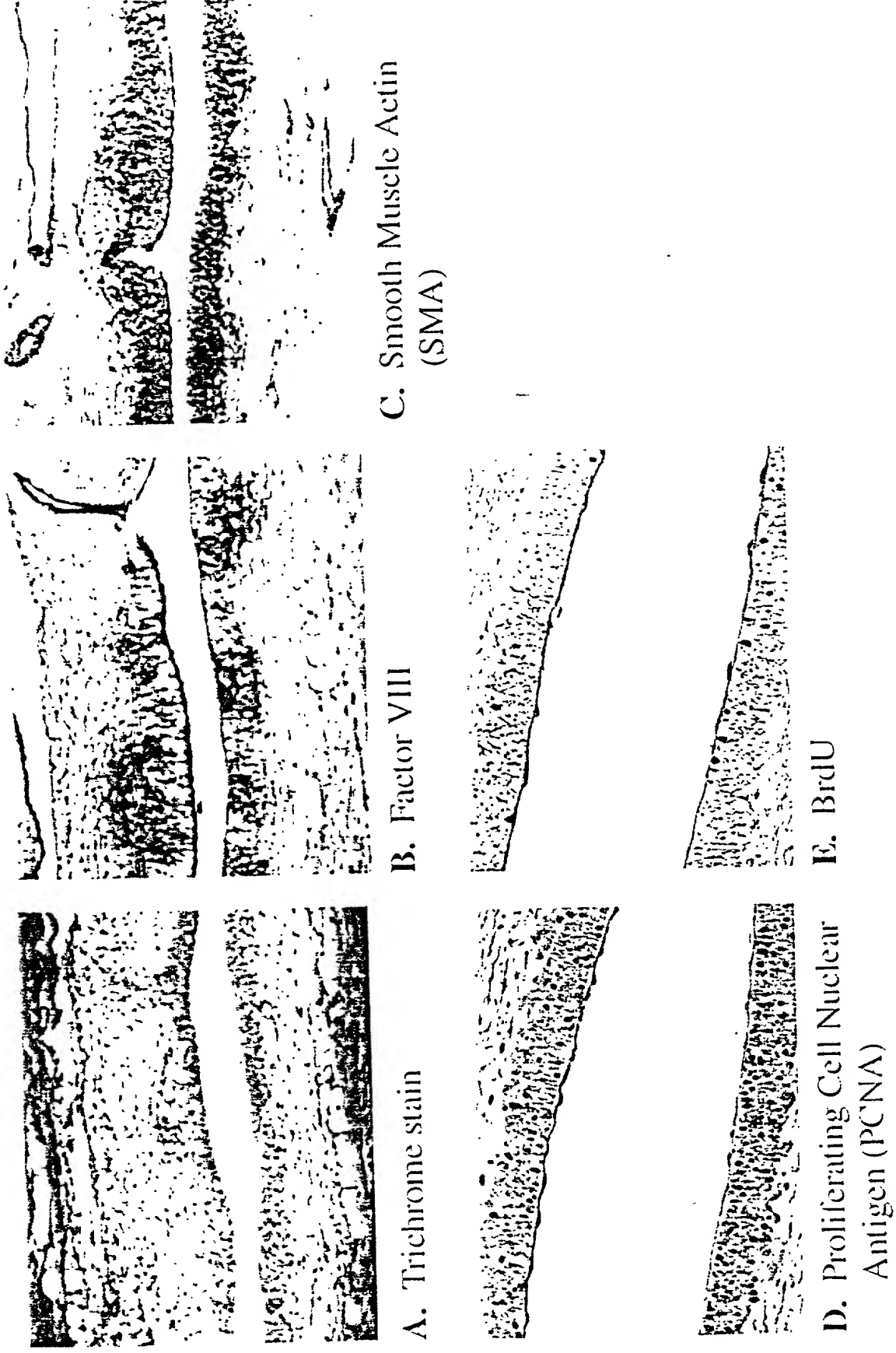
**Figure 20** Quail Chorioallantoic Membrane (CAM) Angiogenesis Assay



**Figure 21** Black and white images of CAM vasculature after growth factor treatment



**Figure 22** Histological sections of blood vessels formed in canine myocardium 2 weeks following BP injection



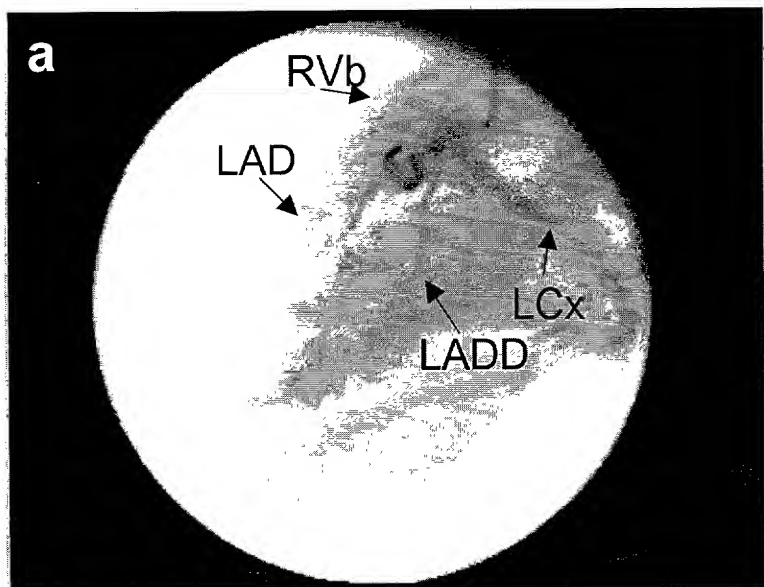


Fig. 23

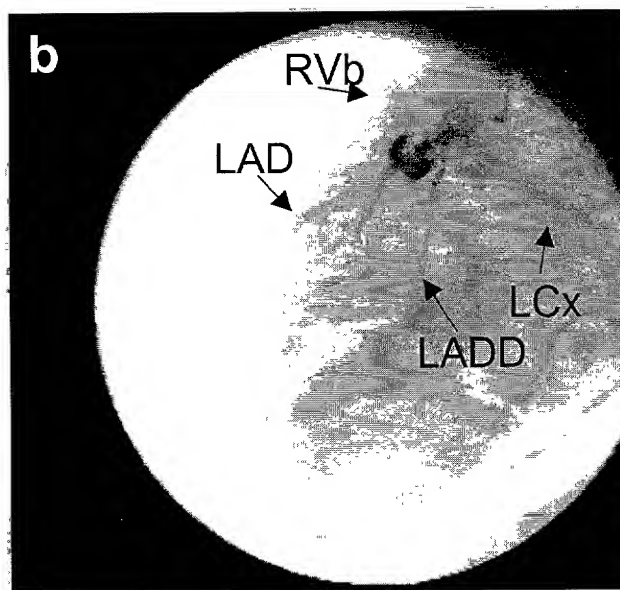


Fig. 24

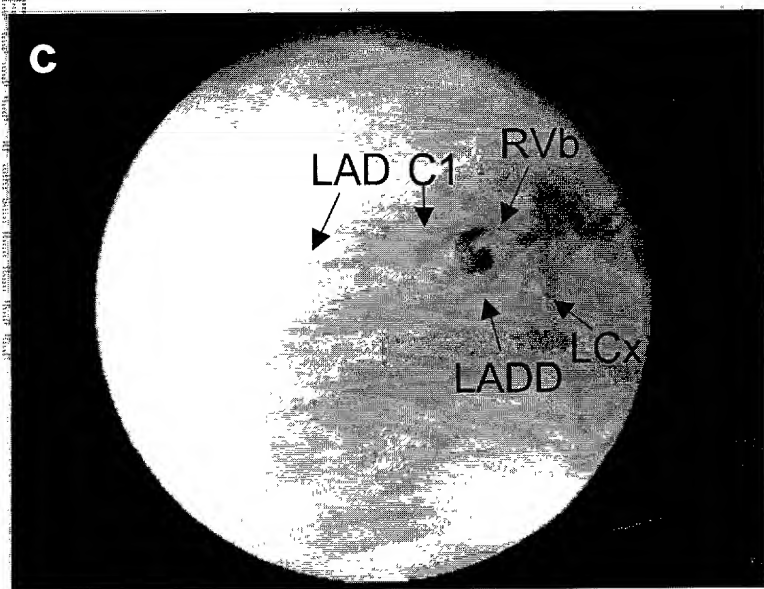


Fig. 25

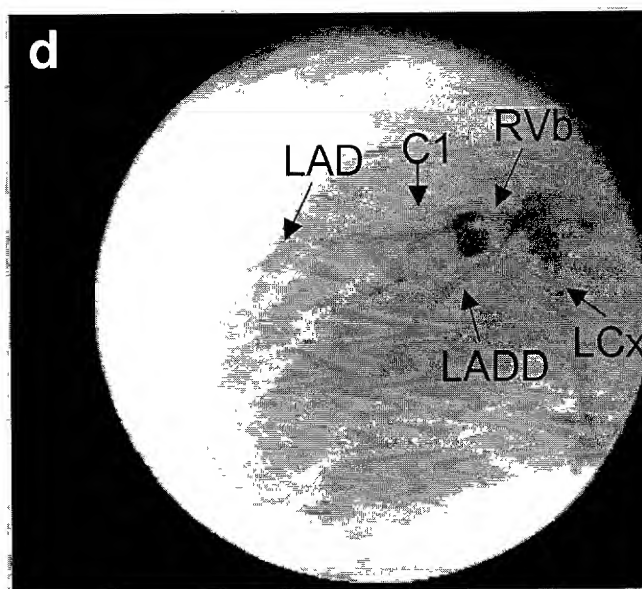


Fig. 26